



SYR-HDAC-5005-C1 replacement 01.ST25.txt
SEQUENCE LISTING

<110> Syrrx, Inc.

<120> HISTONE DEACETYLASE INHIBITORS

<130> SYR-HDAC-5005-C1

<140> US 10/803,575

<141> 2004-03-17

<150> US 60/455,437

<151> 2003-03-17

<150> US 60/531,203

<151> 2003-12-19

<160> 8

<170> PatentIn version 3.2

<210> 1

<211> 513

<212> PRT

<213> Artificial

<220>

<223> Residues 1-482 of HDAC1 and a 6-histidine tag at the N-terminus

<400> 1

Met Ser Tyr Tyr His His His His His Asp Tyr Asp Ile Pro Thr
1 5 10 15

Thr Glu Asn Leu Tyr Phe Gln Gly Ala Met Glu Pro Gly Gly Ser Met
20 25 30

Ala Gln Thr Gln Gly Thr Arg Arg Lys Val Cys Tyr Tyr Tyr Asp Gly
35 40 45

Asp Val Gly Asn Tyr Tyr Tyr Gly Gln Gly His Pro Met Lys Pro His
50 55 60

Arg Ile Arg Met Thr His Asn Leu Leu Leu Asn Tyr Gly Leu Tyr Arg
65 70 75 80

Lys Met Glu Ile Tyr Arg Pro His Lys Ala Asn Ala Glu Glu Met Thr
85 90 95

Lys Tyr His Ser Asp Asp Tyr Ile Lys Phe Leu Arg Ser Ile Arg Pro
100 105 110

Asp Asn Met Ser Glu Tyr Ser Lys Gln Met Gln Arg Phe Asn Val Gly
115 120 125

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Glu Asp Cys Pro Val Phe Asp Gly Leu Phe Glu Phe Cys Gln Leu Ser
 130 135 140
 Thr Gly Gly Ser Val Ala Ser Ala Val Lys Leu Asn Lys Gln Gln Thr
 145 150 155 160
 Asp Ile Ala Val Asn Trp Ala Gly Gly Leu His His Ala Lys Lys Ser
 165 170 175
 Glu Ala Ser Gly Phe Cys Tyr Val Asn Asp Ile Val Leu Ala Ile Leu
 180 185 190
 Glu Leu Leu Lys Tyr His Gln Arg Val Leu Tyr Ile Asp Ile Asp Ile
 195 200 205
 His His Gly Asp Gly Val Glu Glu Ala Phe Tyr Thr Thr Asp Arg Val
 210 215 220
 Met Thr Val Ser Phe His Lys Tyr Gly Glu Tyr Phe Pro Gly Thr Gly
 225 230 235 240
 Asp Leu Arg Asp Ile Gly Ala Gly Lys Gly Lys Tyr Tyr Ala Val Asn
 245 250 255
 Tyr Pro Leu Arg Asp Gly Ile Asp Asp Glu Ser Tyr Glu Ala Ile Phe
 260 265 270
 Lys Pro Val Met Ser Lys Val Met Glu Met Phe Gln Pro Ser Ala Val
 275 280 285
 Val Leu Gln Cys Gly Ser Asp Ser Leu Ser Gly Asp Arg Leu Gly Cys
 290 295 300
 Phe Asn Leu Thr Ile Lys Gly His Ala Lys Cys Val Glu Phe Val Lys
 305 310 315 320
 Ser Phe Asn Leu Pro Met Leu Met Leu Gly Gly Gly Gly Tyr Thr Ile
 325 330 335
 Arg Asn Val Ala Arg Cys Trp Thr Tyr Glu Thr Ala Val Ala Leu Asp
 340 345 350
 Thr Glu Ile Pro Asn Glu Leu Pro Tyr Asn Asp Tyr Phe Glu Tyr Phe
 355 360 365
 Gly Pro Asp Phe Lys Leu His Ile Ser Pro Ser Asn Met Thr Asn Gln
 370 375 380

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Asn Thr Asn Glu Tyr Leu Glu Lys Ile Lys Gln Arg Leu Phe Glu Asn
385 390 395 400

Leu Arg Met Leu Pro His Ala Pro Gly Val Gln Met Gln Ala Ile Pro
405 410 415

Glu Asp Ala Ile Pro Glu Glu Ser Gly Asp Glu Asp Glu Asp Asp Pro
420 425 430

Asp Lys Arg Ile Ser Ile Cys Ser Ser Asp Lys Arg Ile Ala Cys Glu
435 440 445

Glu Glu Phe Ser Asp Ser Glu Glu Glu Gly Glu Gly Gly Arg Lys Asn
450 455 460

Ser Ser Asn Phe Lys Lys Ala Lys Arg Val Lys Thr Glu Asp Glu Lys
465 470 475 480

Glu Lys Asp Pro Glu Glu Lys Lys Glu Val Thr Glu Glu Glu Lys Thr
485 490 495

Lys Glu Glu Lys Pro Glu Ala Lys Gly Val Lys Glu Glu Val Lys Leu
500 505 510

Ala

<210> 2
<211> 1542
<212> DNA
<213> Artificial

<220>
<223> DNA sequence encoding residues 1-482 of HDAC1 and a 6-histidine tag at the N-terminus

<400> 2
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aaagtctgtt actactacga cggggatgtt ggaaattact attatggaca aggccacca 180
atgaagcctc accgaatccg catgactcat aatttgctgc tcaactatgg tctctaccga 240
aaaatggaaa tctatcgccc tcacaaagcc aatgctgagg agatgaccaa gtaccacagc 300
gatgactaca tttaaattctt gcgctccatc cgtccagata acatgtcgga gtacagcaag 360
cagatgcaga gattcaacgt tggtagggac tgtccagtat tcgatggcct gtttgagttc 420
tgtcagttgt ctactggtgg ttctgtggca agtgctgtga aacttaataa gcagcagacg 480

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gacatcgctg tgaattgggc tgggggcctg caccatgcaa agaagtccga ggcattctggc	540
ttctgttacg tcaatgatat cgtcttggcc atcctggaac tgctaaagta tcaccagagg	600
gtgctgtaca ttgacattga tattcaccat ggtgacggcg tggaagaggc cttctacacc	660
acggaccggg tcatgactgt gtcctttcat aagtatggag agtacttccc aggaactggg	720
gacctacggg atatcggggc tggcaaaggc aagtattatg ctgttaacta cccgctccga	780
gacgggattg atgacgagtc ctatgaggcc attttcaagc cggtcatgtc caaagtaatg	840
gagatgttcc agcctagtgc ggtgggtctta cagtgtggct cagactccct atctggggat	900
cggttagggtt gcttcaatct aactatcaaa ggacacgcca agtgtgtgga atttgtcaag	960
agctttaacc tgcctatgct gatgctggga ggcgggtggtt acaccattcg taacgttgcc	1020
cgggtgctgga catatgagac agctgtggcc ctggatacgg agatccctaa tgagcttcca	1080
tacaatgact actttgaata ctttggacca gatttcaagc tccacatcag tccttccaat	1140
atgactaacc agaacacgaa tgagtacctg gagaagatca aacagcgact gtttgagaac	1200
cttagaatgc tgccgcacgc acctggggtc caaatgcagg cgattcctga ggacgccatc	1260
cctgaggaga gtggcgatga ggacgaagac gaccctgaca agcgcattct gatctgctcc	1320
tctgacaaac gaattgcctg tgaggaagag ttctccgatt ctgaagagga gggagagggg	1380
ggccgcaaga actcttccaa cttcaaaaaa gccaagagag tcaaaacaga ggatgaaaaa	1440
gagaaagacc cagaggagaa gaaagaagtc accgaagagg agaaaaccaa ggaggagaag	1500
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<210> 3
 <211> 498
 <212> PRT
 <213> Artificial

<220>
 <223> Residues 1-488 of HDAC2 and a 6-histidine tag at the C-terminus
 <400> 3

Met	Gly	Ser	Met	Ala	Tyr	Ser	Gln	Gly	Gly	Gly	Lys	Lys	Lys	Val	Cys
1				5				10						15	

Tyr	Tyr	Tyr	Asp	Gly	Asp	Ile	Gly	Asn	Tyr	Tyr	Tyr	Gly	Gln	Gly	His
			20				25						30		

Pro	Met	Lys	Pro	His	Arg	Ile	Arg	Met	Thr	His	Asn	Leu	Leu	Leu	Asn
		35					40					45			

Tyr	Gly	Leu	Tyr	Arg	Lys	Met	Glu	Ile	Tyr	Arg	Pro	His	Lys	Ala	Thr
	50					55					60				

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Ala Glu Glu Met Thr Lys Tyr His Ser Asp Glu Tyr Ile Lys Phe Leu
65 70 75 80

Arg Ser Ile Arg Pro Asp Asn Met Ser Glu Tyr Ser Lys Gln Met Gln
85 90 95

Arg Phe Asn Val Gly Glu Asp Cys Pro Val Phe Asp Gly Leu Phe Glu
100 105 110

Phe Cys Gln Leu Ser Thr Gly Gly Ser Val Ala Gly Ala Val Lys Leu
115 120 125

Asn Arg Gln Gln Thr Asp Met Ala Val Asn Trp Ala Gly Gly Leu His
130 135 140

His Ala Lys Lys Ser Glu Ala Ser Gly Phe Cys Tyr Val Asn Asp Ile
145 150 155 160

Val Leu Ala Ile Leu Glu Leu Leu Lys Tyr His Gln Arg Val Leu Tyr
165 170 175

Ile Asp Ile Asp Ile His His Gly Asp Gly Val Glu Glu Ala Phe Tyr
180 185 190

Thr Thr Asp Arg Val Met Thr Val Ser Phe His Lys Tyr Gly Glu Tyr
195 200 205

Phe Pro Gly Thr Gly Asp Leu Arg Asp Ile Gly Ala Gly Lys Gly Lys
210 215 220

Tyr Tyr Ala Val Asn Phe Pro Met Arg Asp Gly Ile Asp Asp Glu Ser
225 230 235 240

Tyr Gly Gln Ile Phe Lys Pro Ile Ile Ser Lys Val Met Glu Met Tyr
245 250 255

Gln Pro Ser Ala Val Val Leu Gln Cys Gly Ala Asp Ser Leu Ser Gly
260 265 270

Asp Arg Leu Gly Cys Phe Asn Leu Thr Val Lys Gly His Ala Lys Cys
275 280 285

Val Glu Val Val Lys Thr Phe Asn Leu Pro Leu Leu Met Leu Gly Gly
290 295 300

Gly Gly Tyr Thr Ile Arg Asn Val Ala Arg Cys Trp Thr Tyr Glu Thr
305 310 315 320

SYR-HDAC-5005-C1 replacement 01.ST25.txt

Ala Val Ala Leu Asp Cys Glu Ile Pro Asn Glu Leu Pro Tyr Asn Asp
325 330 335

Tyr Phe Glu Tyr Phe Gly Pro Asp Phe Lys Leu His Ile Ser Pro Ser
340 345 350

Asn Met Thr Asn Gln Asn Thr Pro Glu Tyr Met Glu Lys Ile Lys Gln
355 360 365

Arg Leu Phe Glu Asn Leu Arg Met Leu Pro His Ala Pro Gly Val Gln
370 375 380

Met Gln Ala Ile Pro Glu Asp Ala Val His Glu Asp Ser Gly Asp Glu
385 390 395 400

Asp Gly Glu Asp Pro Asp Lys Arg Ile Ser Ile Arg Ala Ser Asp Lys
405 410 415

Arg Ile Ala Cys Asp Glu Glu Phe Ser Asp Ser Glu Asp Glu Gly Glu
420 425 430

Gly Gly Arg Arg Asn Val Ala Asp His Lys Lys Gly Ala Lys Lys Ala
435 440 445

Arg Ile Glu Glu Asp Lys Lys Glu Thr Glu Asp Lys Lys Thr Asp Val
450 455 460

Lys Glu Glu Asp Lys Ser Lys Asp Asn Ser Gly Glu Lys Thr Asp Thr
465 470 475 480

Lys Gly Thr Lys Ser Glu Gln Leu Ser Asn Pro Gly His His His His
485 490 495

His His

<210> 4
<211> 1497
<212> DNA
<213> Artificial

<220>
<223> DNA sequence encoding residues 1-488 of HDAC2 and a 6-histidine tag at the C-terminus

<400> 4
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ggtgatattg gaaattatta ttatggacag ggtcatccca tgaagcctca tagaatccgc 120

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atgacccata acttgctgtt aaattatggc ttatacagaa aaatggaaat atataggccc 180
cataaagcca ctgccgaaga aatgacaaaa tatcacagtg atgagtatat caaatttcta 240
cgggtcaataa gaccagataa catgtctgag tatagtaagc agatgcagag atttaatgtt 300
ggagaagatt gtccagtgtt tgatggactc tttgagtttt gtcagctctc aactggcggg 360
tcagttgctg gagctgtgaa gttaaaccga caacagactg atatggctgt taattgggct 420
ggaggattac atcatgctaa gaaatcagaa gcatcaggat tctgttacgt taatgatatt 480
gtgcttgcca tccttgaatt actaaagtat catcagagag tcttatatat tgatatagat 540
attcatcatg gtgatgggtg tgaagaagct ttttatacaa cagatcgtgt aatgacggta 600
tcattccata aatatgggga atactttcct ggcacaggag acttgaggga tattggtgct 660
ggaaaaggca aatactatgc tgtcaatttt ccaatgagag atggtataga tgatgagtca 720
tatgggcaga tatttaagcc tattatctca aaggatgatg agatgtatca acctagtgt 780
gtggtattac agtgtggtgc agactcatta tctggtgata gactggggtt tttcaatcta 840
acagtcaaag gtcattgctaa atgtgtagaa gttgtaaaaa cttttaactt accattactg 900
atgcttggag gaggtggcta cacaatccgt aatgttgctc gatgttggac atatgagact 960
gcagttgccc ttgattgtga gattcccaat gagttgccat ataatgatta ctttgagtat 1020
tttgaccag acttcaaact gcatattagt ctttcaaaca tgacaaacca gaacactcca 1080
gaatatatgg aaaagataaa acagcgtttg tttgaaaatt tgcgcatgtt acctcatgca 1140
cctggtgtcc agatgcaagc tattccagaa gatgctgttc atgaagacag tggagatgaa 1200
gatggagaag atccagacaa gagaatttct attcgagcat cagacaagcg gatagcttgt 1260
gatgaagaat tctcagattc tgaggatgaa ggagaaggag gtcgaagaaa tgtggctgat 1320
cataagaaag gagcaaagaa agctagaatt gaagaagata agaaagaaac agaggacaaa 1380
aaaacagacg ttaaggaaga agataaatcc aaggacaaca gtggtgaaaa aacagatacc 1440
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<210> 5
 <211> 782
 <212> PRT
 <213> Artificial

<220>
 <223> Residues 73-845 of HDAC6 and a 6-histidine tag at the C-terminus

<400> 5

Met Pro Gly Met Asp Leu Asn Leu Glu Ala Glu Ala Leu Ala Gly Thr
 1 5 10 15

Gly Leu Val Leu Asp Glu Gln Leu Asn Glu Phe His Cys Leu Trp Asp
 20 25 30

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Asp Ser Phe Pro Glu Gly Pro Glu Arg Leu His Ala Ile Lys Glu Gln
 35 40 45
 Leu Ile Gln Glu Gly Leu Leu Asp Arg Cys Val Ser Phe Gln Ala Arg
 50 55 60
 Phe Ala Glu Lys Glu Glu Leu Met Leu Val His Ser Leu Glu Tyr Ile
 65 70 75 80
 Asp Leu Met Glu Thr Thr Gln Tyr Met Asn Glu Gly Glu Leu Arg Val
 85 90 95
 Leu Ala Asp Thr Tyr Asp Ser Val Tyr Leu His Pro Asn Ser Tyr Ser
 100 105 110
 Cys Ala Cys Leu Ala Ser Gly Ser Val Leu Arg Leu Val Asp Ala Val
 115 120 125
 Leu Gly Ala Glu Ile Arg Asn Gly Met Ala Ile Ile Arg Pro Pro Gly
 130 135 140
 His His Ala Gln His Ser Leu Met Asp Gly Tyr Cys Met Phe Asn His
 145 150 155 160
 Val Ala Val Ala Ala Arg Tyr Ala Gln Gln Lys His Arg Ile Arg Arg
 165 170 175
 Val Leu Ile Val Asp Trp Asp Val His His Gly Gln Gly Thr Gln Phe
 180 185 190
 Thr Phe Asp Gln Asp Pro Ser Val Leu Tyr Phe Ser Ile His Arg Tyr
 195 200 205
 Glu Gln Gly Arg Phe Trp Pro His Leu Lys Ala Ser Asn Trp Ser Thr
 210 215 220
 Thr Gly Phe Gly Gln Gly Gln Gly Tyr Thr Ile Asn Val Pro Trp Asn
 225 230 235 240
 Gln Val Gly Met Arg Asp Ala Asp Tyr Ile Ala Ala Phe Leu His Val
 245 250 255
 Leu Leu Pro Val Ala Leu Glu Phe Gln Pro Gln Leu Val Leu Val Ala
 260 265 270
 Ala Gly Phe Asp Ala Leu Gln Gly Asp Pro Lys Gly Glu Met Ala Ala
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275

280

285

Thr Pro Ala Gly Phe Ala Gln Leu Thr His Leu Leu Met Gly Leu Ala
290 295 300

Gly Gly Lys Leu Ile Leu Ser Leu Glu Gly Gly Tyr Asn Leu Arg Ala
305 310 315 320

Leu Ala Glu Gly Val Ser Ala Ser Leu His Thr Leu Leu Gly Asp Pro
325 330 335

Cys Pro Met Leu Glu Ser Pro Gly Ala Pro Cys Arg Ser Ala Gln Ala
340 345 350

Ser Val Ser Cys Ala Leu Glu Ala Leu Glu Pro Phe Trp Glu Val Leu
355 360 365

Val Arg Ser Thr Glu Thr Val Glu Arg Asp Asn Met Glu Glu Asp Asn
370 375 380

Val Glu Glu Ser Glu Glu Glu Gly Pro Trp Glu Pro Pro Val Leu Pro
385 390 395 400

Ile Leu Thr Trp Pro Val Leu Gln Ser Arg Thr Gly Leu Val Tyr Asp
405 410 415

Gln Asn Met Met Asn His Cys Asn Leu Trp Asp Ser His His Pro Glu
420 425 430

Val Pro Gln Arg Ile Leu Arg Ile Met Cys Arg Leu Glu Glu Leu Gly
435 440 445

Leu Ala Gly Arg Cys Leu Thr Leu Thr Pro Arg Pro Ala Thr Glu Ala
450 455 460

Glu Leu Leu Thr Cys His Ser Ala Glu Tyr Val Gly His Leu Arg Ala
465 470 475 480

Thr Glu Lys Met Lys Thr Arg Glu Leu His Arg Glu Ser Ser Asn Phe
485 490 495

Asp Ser Ile Tyr Ile Cys Pro Ser Thr Phe Ala Cys Ala Gln Leu Ala
500 505 510

Thr Gly Ala Ala Cys Arg Leu Val Glu Ala Val Leu Ser Gly Glu Val
515 520 525

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Leu Asn Gly Ala Ala Val Val Arg Pro Pro Gly His His Ala Glu Gln
530 535 540

Asp Ala Ala Cys Gly Phe Cys Phe Phe Asn Ser Val Ala Val Ala Ala
545 550 555 560

Arg His Ala Gln Thr Ile Ser Gly His Ala Leu Arg Ile Leu Ile Val
565 570 575

Asp Trp Asp Val His His Gly Asn Gly Thr Gln His Met Phe Glu Asp
580 585 590

Asp Pro Ser Val Leu Tyr Val Ser Leu His Arg Tyr Asp His Gly Thr
595 600 605

Phe Phe Pro Met Gly Asp Glu Gly Ala Ser Ser Gln Ile Gly Arg Ala
610 615 620

Ala Gly Thr Gly Phe Thr Val Asn Val Ala Trp Asn Gly Pro Arg Met
625 630 635 640

Gly Asp Ala Asp Tyr Leu Ala Ala Trp His Arg Leu Val Leu Pro Ile
645 650 655

Ala Tyr Glu Phe Asn Pro Glu Leu Val Leu Val Ser Ala Gly Phe Asp
660 665 670

Ala Ala Arg Gly Asp Pro Leu Gly Gly Cys Gln Val Ser Pro Glu Gly
675 680 685

Tyr Ala His Leu Thr His Leu Leu Met Gly Leu Ala Ser Gly Arg Ile
690 695 700

Ile Leu Ile Leu Glu Gly Gly Tyr Asn Leu Thr Ser Ile Ser Glu Ser
705 710 715 720

Met Ala Ala Cys Thr Arg Ser Leu Leu Gly Asp Pro Pro Pro Leu Leu
725 730 735

Thr Leu Pro Arg Pro Pro Leu Ser Gly Ala Leu Ala Ser Ile Thr Glu
740 745 750

Thr Ile Gln Val His Arg Arg Tyr Trp Arg Ser Leu Arg Val Met Lys
755 760 765

Val Glu Asp Arg Glu Gly Pro Gly His His His His His His
770 775 780

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<210> 6
 <211> 2349
 <212> DNA
 <213> Artificial

<220>
 <223> DNA encoding residues 73-845 of HDAC6 and a 6-histidine tag at the C-terminus

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 cggctccatg ccatcaagga gcaactgac caggagggcc tcctagatcg ctgcgtgtcc 180
 ttccaggccc gggttgctga aaaggaagag ctgatgttgg ttcacagcct agaatatatt 240
 gatctgatgg aaacaacca gtacatgaat gagggagaac tccgtgtcct agcagacacc 300
 tacgactcag ttatctgca tccgaactca tactcctgtg cctgcctggc ctcaggctct 360
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 gattgggatg tgcaccacgg tcaaggaaca cagttcacct tcgaccagga cccagtgctc 600
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 aactggtcca ccacaggttt cggccaaggc caaggatata ccatcaatgt gccttggaac 720
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 aatcactgca acttggtgga cagccaccac cctgaggtac cccagcgcac cttgcggatc 1320
 atgtgccgtc tggaggagct gggccttgcc gggcgctgcc tcacctgac accgcgccct 1380
 gccacagagg ctgagctgct cacctgtcac agtgctgagt acgtgggtca tctccgggcc 1440
 acagagaaaa tgaaaacccg ggagctgcac cgtgagagtt ccaactttga ctccatctat 1500
 atctgccccca gtaccttcgc ctgtgcacag cttgccactg gcgctgcctg ccgcctggtg 1560

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gaggctgtgc tctcaggaga ggttctgaat ggtgctgctg tggcgctcc cccaggacac	1620
cacgcagagc aggatgcagc ttgcggtttt tgctttttca actctgtggc tgtggctgct	1680
cgccatgccc agactatcag tgggcatgcc ctacggatcc tgattgtgga ttgggatgtc	1740
caccacggta atggaactca gcacatgttt gaggatgacc ccagtgtgct atatgtgtcc	1800
ctgcaccgct atgatcatgg caccttcttc cccatggggg atgaggggtgc cagcagccag	1860
atcggccggg ctgcgggcac aggcttcacc gtcaacgtgg catggaacgg gccccgcatg	1920
ggtgatgctg actacctagc tgcctggcat cgcctggtgc ttcccattgc ctacgagttt	1980
aaccacagaac tgggtgctggt ctcagctggc tttgatgctg cacgggggga tccgctgggg	2040
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atggctgcct gcactcgctc cctccttgga gaccaccac ccctgctgac cctgccacgg	2220
ccccactat caggggccct ggcctcaatc actgagacca tccaagtcca tcgcagatac	2280
tggcgagct tacgggtcat gaaggtagaa gacagagaag gacccgggca tcaccatcac	2340
catcactaa	2349

<210> 7
 <211> 385
 <212> PRT
 <213> Artificial

<220>
 <223> Residues 1-377 of HDAC8 and a 6-histidine tag at the N-terminus
 <400> 7

Met	His	His	His	His	His	His	Pro	Met	Glu	Glu	Pro	Glu	Glu	Pro	Ala
1				5					10					15	

Asp	Ser	Gly	Gln	Ser	Leu	Val	Pro	Val	Tyr	Ile	Tyr	Ser	Pro	Glu	Tyr
		20						25					30		

Val	Ser	Met	Cys	Asp	Ser	Leu	Ala	Lys	Ile	Pro	Lys	Arg	Ala	Ser	Met
		35					40					45			

Val	His	Ser	Leu	Ile	Glu	Ala	Tyr	Ala	Leu	His	Lys	Gln	Met	Arg	Ile
	50					55					60				

Val	Lys	Pro	Lys	Val	Ala	Ser	Met	Glu	Glu	Met	Ala	Ala	Phe	His	Thr
65					70					75					80

Asp	Ala	Tyr	Leu	Gln	His	Leu	Gln	Lys	Val	Ser	Gln	Glu	Gly	Asp	Asp
			85						90					95	

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Asp His Pro Asp Ser Ile Glu Tyr Gly Leu Gly Tyr Asp Cys Pro Ala
100 105 110

Thr Glu Gly Ile Phe Asp Tyr Ala Ala Ala Ile Gly Gly Ala Thr Ile
115 120 125

Thr Ala Ala Gln Cys Leu Ile Asp Gly Met Cys Lys Val Ala Ile Asn
130 135 140

Trp Ser Gly Gly Trp His His Ala Lys Lys Asp Glu Ala Ser Gly Phe
145 150 155 160

Cys Tyr Leu Asn Asp Ala Val Leu Gly Ile Leu Arg Leu Arg Arg Lys
165 170 175

Phe Glu Arg Ile Leu Tyr Val Asp Leu Asp Leu His His Gly Asp Gly
180 185 190

Val Glu Asp Ala Phe Ser Phe Thr Ser Lys Val Met Thr Val Ser Leu
195 200 205

His Lys Phe Ser Pro Gly Phe Phe Pro Gly Thr Gly Asp Val Ser Asp
210 215 220

Val Gly Leu Gly Lys Gly Arg Tyr Tyr Ser Val Asn Val Pro Ile Gln
225 230 235 240

Asp Gly Ile Gln Asp Glu Lys Tyr Tyr Gln Ile Cys Glu Ser Val Leu
245 250 255

Lys Glu Val Tyr Gln Ala Phe Asn Pro Lys Ala Val Val Leu Gln Leu
260 265 270

Gly Ala Asp Thr Ile Ala Gly Asp Pro Met Cys Ser Phe Asn Met Thr
275 280 285

Pro Val Gly Ile Gly Lys Cys Leu Lys Tyr Ile Leu Gln Trp Gln Leu
290 295 300

Ala Thr Leu Ile Leu Gly Gly Gly Gly Tyr Asn Leu Ala Asn Thr Ala
305 310 315 320

Arg Cys Trp Thr Tyr Leu Thr Gly Val Ile Leu Gly Lys Thr Leu Ser
325 330 335

Ser Glu Ile Pro Asp His Glu Phe Phe Thr Ala Tyr Gly Pro Asp Tyr
340 345 350

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Val Leu Glu Ile Thr Pro Ser Cys Arg Pro Asp Arg Asn Glu Pro His
355 360 365

Arg Ile Gln Gln Ile Leu Asn Tyr Ile Lys Gly Asn Leu Lys His Val
370 375 380

Val
385

<210> 8
<211> 1158
<212> DNA
<213> Artificial

<220>
<223> DNA encoding residues 1-377 of HDAC8 and a 6-histidine tag at the N-terminus

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